



**EMBL-EBI**  
European Bioinformatics Institute

Get  for  Go ? Site search

[EBI Home](#)
[About EBI](#)
[Research](#)
[Services](#)
[Toolbox](#)
[Databases](#)
[Downloads](#)
[Submit](#)

DATABASE BROWSING

## EBI Dbfetch

```

ID  HPCCORBEA  standard; genomic RNA; VRL; 447 BP.
XX
AC  L39317;
XX
SV  L39317.1
XX
DT  30-JUN-1995 (Rel. 44, Created)
DT  21-OCT-2001 (Rel. 69, Last updated, Version 4)
XX
DE  Hepatitis C virus type 3 clone NL96 precursor protein gene, partial cds.
XX
KW  .
XX
OS  Hepatitis C virus type 3
OC  Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
OC  Hepacivirus.
XX
RN  [1]
RP  1-447
RX  MEDLINE; 95052487.
RX  PUBMED; 7525693.
RA  van Doorn L.J., Kleter B., Stuyver L., Maertens G., Brouwer H., Schalm S.,
RA  Heijntink R., Quint W.;
RT  "Analysis of hepatitis C virus genotypes by a line probe assay and
RT  correlation with antibody profiles";
RL  J. Hepatol. 21(1):122-129(1994).
XX
RN  [2]
RP  1-447
RX  MEDLINE; 97201609.
RX  PUBMED; 9049395.
RA  van Doorn L.J., Kleter G.E., Stuyver L., Maertens G., Brouwer J.T.,
RA  Schalm S.W., Heijntink R.A., Quint W.G.V.;
RT  "Sequence analysis of hepatitis C virus genotypes 1 to 5 reveals multiple
RT  novel subtypes in the Benelux countries";
RL  J. Gen. Virol. 76(Pt 7):1871-1876(1995).
XX
FH  Key          Location/Qualifiers
FH
FT  source      1..447
FT                /db_xref="taxon:40363"
FT                /mol_type="genomic RNA"
FT                /note="genotype: 3"
FT                /organism="Hepatitis C virus type 3"
FT                /clone="NL96"
FT  mat_peptide 1..96
FT                /note="putative"
FT                /product="core protein"
FT  CDS         <1..>447
FT                /codon_start=1
FT                /db_xref="GOA:Q81286"
FT                /db_xref="InterPro:IPR002519"
FT                /db_xref="InterPro:IPR002521"
FT                /db_xref="UniProt/TreMBL:Q81286"
FT                /product="precursor protein"
FT                /protein_id="AAA67840.1"

```

```

FT      /translation="DGINFATGNLPGCSFSIFLLALFSCLLTPTAGLEYRNASGLYMT
FT      NDCSNGSIVYEAGDIILHLPGCVPCVRSNGT SRCWIPVSPTVAVKSPCAATASLRTHVD
FT      MMVGAATLCSALYVGDLGALFLVGQGF SWRHRQHWTVQDCNC SI"
FT      mat peptide      97..447
FT      /note="putative"
FT      /product="el protein"
XX
SQ      Sequence 447 BP; 82 A; 130 C; 114 G; 118 T; 3 other;
      gacggaatta atttcgcaac agggaattta cctggttgct ctttctctat cttccttctg      60
      gctttgttct catgcttgct tacacccaca gccgggctgg agtaccgtaa tgcctccgga      120
      ctctacatgg taactaacga ctgcagtaac ggtagtatcg tgtatgaggc cggggatatt      180
      atcctccact tacctggctg tgtcccctgc gtacgctctg gcaatacatc aagatgctgg      240
      atccctgtga gcccyaaccg cgccgtgaag tcgccctgcg ccgccaccgc ctctctccgc      300
      acgcacgtgg atatgatggg gggrgcggcc accctatgct cagctctcta cgtaggagac      360
      ctttgtggag cgctatttct tgytyggcag gggttctcat ggagacatcg ccagcattgg      420
      actgtccagg actgcaactg ttccatc      447
//

```